

Figure S1. 150 mmol/L EtOH incubation induces steatosis in L02 cells and steatosis was ameliorated by SR8278 treatment or Rev-erbashRNA transfection in EtOHtreated L-02 cells. (A) (B) Oil Red O staining and the Content of TG analysis in Control and EtOH group (Scale bar =100 µm). (C) (D) qRT-PCR and Western blot analysis of Pparα and Srebp1c in Control and EtOH group. EtOH-treated L-02 cells were treated with or without 10 µM SR8278 for 24 h. (E) (F) Oil Red O staining and TG content in EtOH-treated L-02 cells treated with SR8278 (Scale bar =100 μm). (G) Western blot analysis of Ppara and Srebp1c in EtOH-treated L-02 cells after SR8278 treatment. (H) qRT-PCR analysis of Rev-erba in EtOH-treated L-02 cells transfected with ReverbashRNA. (n=3). Bar represents the mean \pm S.E.M.. Significance * P < 0.05, ** P<0.01 vs. Control group. ${}^{\#}P < 0.05$, ${}^{\#\#}P < 0.01$ vs. EtOH group.

Ppara

Srebp1c

EtOH

SR8278

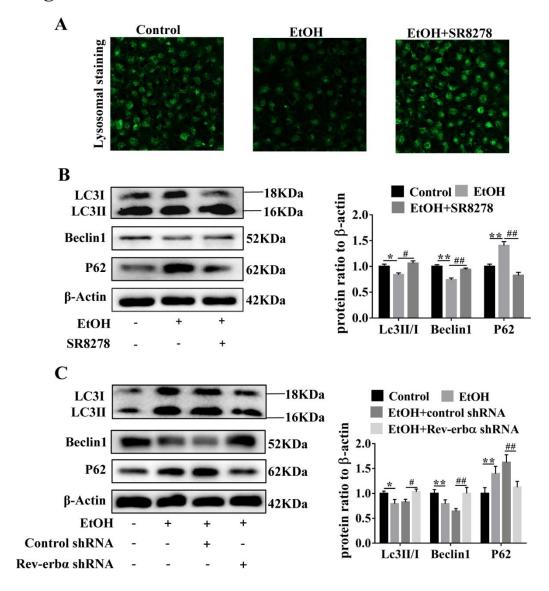


Figure S2. Autophagy activity was improved by SR8278 treatment or ReverbαshRNA transfection in EtOH-treated L-02 cells. (A) Lysosome staining with Lyso Tracter Green DND-26 in EtOH-treated L-02 cells treated with SR8278 (Scale bars=40 μm). (B) Western blot analysis of Lc3II/I, Beclin1 and P62 in EtOH-treated L-02 cells treated with SR8278. (C) Western blot analysis of Lc3II/I, Beclin1 and P62 in EtOH-treated L-02 cells in EtOH-treated L-02 cells transfected with Rev-erbα shRNA (n \geq 3). Bar represents the mean \pm S.E.M.. Significance * P < 0.05, ** P < 0.01 vs. Control group. # P < 0.05, ## P < 0.01 vs. Control shRNA group.

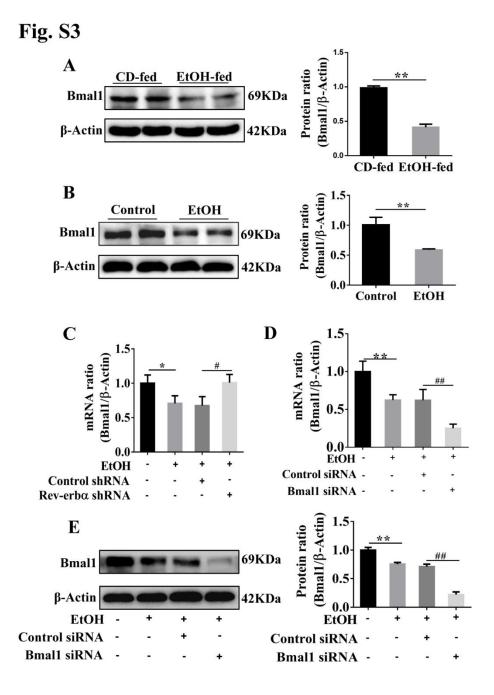


Figure S3. Bmal1 was up-regulated in EtOH-treated L-02 cells and was regulated by Rev-erbα. (A) Western blot analysis of Bmal1 in the liver of EtOH-fed mice (n≥6). (B) Western blot analysis of Bmal1 in EtOH-treated L-02 cells (n ≥3). Bar represents the mean \pm S.E.M.. Significance * P < 0.05, ** P < 0.01 vs. EtOH-fed group or EtOH group. (C) qRT-PCR analysis of Bmal1 in EtOH-treated L-02 cells transfected with Rev-erbα shRNA (n ≥3). Bar represents the mean \pm S.E.M.. Significance * P < 0.05, ** P < 0.01 vs. Control group, # P < 0.05, ## P < 0.01 vs. Control shRNA group. (D) (E) qRT-PCR and western blot analysis of Bmal1 in EtOH-treated L-02 cells transfected with Bmal1 siRNA (n ≥3). Bar represents the mean \pm S.E.M.. Significance * P < 0.05, ** P < 0.05, ** P < 0.01 vs. Control group, # P < 0.05, ## P < 0.05, ## P < 0.01 vs. Control siRNA group.